



# Effects of S-Abscissic Acid (S-ABA) on Seed Germination, Seedling Growth, and *Asr1* Gene Expression Under Drought Stress in Maize

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## Abstract

The objective of this study was to evaluate the ability of the phytohormone S-abscissic acid (S-ABA) to protect maize seedlings grown under drought stress and to measure their increased drought tolerance. The maize hybrids ‘Zhengdan 958’ (ZD958; drought tolerant) and ‘Xundan 20’ (XD20; drought sensitive) were treated with nutrient solutions of different concentrations (1, 2, 4, 8, and 10 mg/kg) of S-ABA under polyethylene glycol (PEG, 15% w/v, MW 6000) simulated drought stress. Optimal concentrations of S-ABA were designed to be sprayed onto the leaves of seedlings, and their effect on endogenous ABA, malondialdehyde (MDA), osmotic substances, antioxidant enzyme activities, and *Asr1* gene expression in seedlings were studied. Results indicated that, under drought stress, S-ABA treatment significantly improved maize seed germination rate (GR), germination energy (GE), and seedling biomass ( $p < 0.05$ ). After spraying 4 mg/kg S-ABA onto leaves, the endogenous hormone ABA, osmotic substances, antioxidant enzyme activities, and expressive quantity of the *Asr1* gene were extended and MDA content dropped significantly ( $p < 0.05$ ). Moreover, ZD 958 endogenous ABA content, osmotic substances content, antioxidant enzyme activity and *Asr1* gene expressive quantity were higher than that of XD 20 ( $p < 0.05$ ). In conclusion, S-ABA treatment increased the content of endogenous ABA, induced an increase in antioxidant enzyme activity and *Asr1* gene expression level, reduced the oxidative damage caused by drought to maize leaves, and improved the adaptability of maize seedlings to withstand drought stress. The promoting effect of S-ABA on the drought-tolerant variety ZD 958 was more obvious ( $p < 0.05$ ). These results serve as a reference for the use of S-ABA in mitigating drought stress in maize.

**Keywords** S-ABA · Drought · Maize · Endogenous hormones · *Asr1* · Antioxidant enzyme

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Chentao Yao and Fengwen Zhang are the first authors. They contributed equally to the writing of this article.

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## Introduction

Drought is an important topic that urgently needs to be addressed in the light of rising global temperatures. Only four-fifths of global arable land has natural precipitation as its main source of water (Turner 2004) and of this, more than 30% falls in arid or semi-arid areas, spread over many countries and regions (Zhao et al. 2007). It is one of the major constraints to the cultivation of maize (*Zea mays* L.) crops.

In China, arid and semi-arid areas account for about half of the country’s land area and these face the threat of periodic drought (Leng et al. 2015). Maize is one of the most important cereals grown globally. Normally, the crop needs 500–800 mm of water during its life cycle (80–110 days). The consequences of drought stress in crop production systems are perhaps more deleterious than abiotic stress under changing climatic scenarios, which seriously restrict maize yield (Badu-Apraku et al. 2013). The Huang-Huai-hai area

is an important region for the planting of summer maize, and here the occurrence of drought stress during the growing period may hamper water-use efficiency leading to significant yield losses.

Throughout evolution, plants have established complicated self-defense mechanisms, including antioxidant defense systems such as superoxide dismutase (SOD, EC 1.15.1.1), catalase (CAT, EC 1.11.1.6), peroxidase (POD, EC 1.11.1.7), ascorbate peroxidase (APX, EC 1.11.1.11), polyphenol oxidase (PPO, EC 1.10.3.2), and phenylalanine ammonia lyase (PAL, EC 4.3.1.5), and non-antioxidant defense systems such as Pro, soluble protein, and so on (Ge et al. 2006; Hawrylak-Nowak et al. 2018; Tan et al. 2011). Many studies have demonstrated a correlation between the activity of these protective enzymes and water stress (Srivalli et al. 2010). At the same time, drought stress induces the expression of related genes in plants. Abscisic acid-stress-ripening-induced (ASR) proteins are a group of plant-specific proteins that are hydrophilic with a low molecular weight and are considered to be transcription factors that were first discovered in tomato plants. Genes of the *Asr* family were characterized by their strong induction under abiotic stress and ABA signaling. Their members harbor the ABA/WDS (abscisic acid/water deficit stress) domain as a common denominator (Virilouvet et al. 2011). The function of the *Asr* gene is mainly to regulate plant growth and development, senescence, fruit ripening, glucose metabolism, and response to stress (Dominguez et al. 2013; Iusem et al. 1993; Yang et al. 2008). Under water stress, *Asr1* overexpression in maize plants increased yield, which suggests that *Asr1* participates in plant growth through the regulation of kind of metabolites (Virilouvet et al. 2011). In a wide variety of species including lily, maize, rice, tobacco, and banana, the expression of *Asr1* can be induced by ABA, thereby enhancing crop drought tolerance (Hu et al. 2013; Joo et al. 2013; Virilouvet et al. 2011; Yang et al. 2005). S-Abscisic acid (S-ABA; scientific name: abscisic acid) is an endogenous plant hormone with important physiological activities. Recent research has found that S-ABA is an “anti-hard environments inducing factor” that can induce a resistance in plants against harsh environments, such as drought and other stressful situations.

To increase crop yields in water-deficit areas, it is crucially important to induce drought tolerance in plants and several agronomical and physiological practices have been developed to achieve this goal. The effects of drought stress on growth and development, eco-physiological characteristics, and the expression levels of various genes in maize have been studied by many scholars in the recent years. Du et al. (2013) indicate that the application of ABA can reduce the rate of transpiration loss and reduce production. The use of S-ABA hormone before heading and flowering to alleviate heat damage to rice under drought conditions is reported

by Gao et al. (2015). Li et al. (2010) show that the application of ABA can increase antioxidant enzyme activity in sugarcane leaves, reduce the content of  $H_2O_2$  and MDA, reduce membrane lipid peroxidation, and then enhance the antioxidant defense system of sugarcane and improve drought resistance. Ghosh et al. (2015) show that the ASR gene is present in the potato and its expression is induced by the stress of dehydration and abscisic acid. Sugiharto et al. (2002) report the cloning of the ASR gene in sugarcane, and their results show that the expression of this gene is induced by ABA treatment. The ability of S-ABA to increase tolerance to drought stress has been reported in rice, cotton, and wheat (Li et al. 2017; Travaglia et al. 2007; Yan et al. 2014). In spite of this, there have been relatively few studies on the regulation of drought resistance of maize by S-ABA and very little is known about the interaction between S-ABA and drought-tolerant maize genotypes or drought-sensitive maize genotypes.

In this context, the objective of this study was to evaluate the possible roles of the exogenous application of ABA for increasing drought tolerance in drought-tolerant and drought-sensitive maize hybrids based on *Asr1* gene-expressive quantity and physiological changes such as endogenous ABA and antioxidant enzyme activity.

## Materials and Methods

The concentrated (98%) S-ABA pesticide used in this study was obtained from Sichuan Dragon Python Fusheng Technology Co. Ltd. Aliquots of 0.5102 g of 98% S-ABA were dissolved in acetone to a constant volume of 50 ml in a volumetric flask to obtain the mother liquor A at a concentration of 10,000 mg/kg. Volumes of 1 ml of mother liquor A were placed into 100 ml volumetric flasks using a pipette gun (Eppendorf), and the volume was kept constant by the addition of 0.1% Tween 80. Finally, this solution was diluted to obtain 1, 2, 4, 8, 10 mg/kg of S-ABA.

## Experimental Setup and Stress Conditions

Maize (*Zea mays* L.) seeds were surface-sterilized with 10% NaClO for 10 min and rinsed thoroughly with distilled water then placed on neutral filter paper to dry. Under the conditions simulated in 15% PEG-6000 experiments, 1, 2, 4, 8, and 10 mg/kg (active principle) concentrations of the commercial raw pesticide S-ABA were prepared in which to soak the seeds. After soaking for 10 h, the seeds were removed and placed into a 15 cm Petri dish with two layers of filter paper, setting blank controls for drought and using distilled water; these were named CK2 and CK1, respectively. The seeds were grown in a 28 °C constant temperature incubator. For this purpose, an experiment was carried out with three

replications and 30 seeds in each petri dish. The number of seeds that germinated from the third day was counted, and the germination energy (GE) and germination rate (GR) were calculated. At the end of the eighth day of the germination test, ten plants were sampled randomly to measure the root length, shoot length, and the dry and fresh weight of roots and shoots. Before recording the plant dry biomass, harvested plants were cut into pieces and placed in an oven at 80 °C until a constant weight was achieved.

GE = The number naturally germinated within 4 days/test seeds  $\times$  100%

GR = The number naturally germinated within 7 days/test seeds  $\times$  100%

To study the physiological effects of S-ABA on maize seedlings under drought conditions, a sand culture was incubated in a constant-temperature incubator at 28 °C, and Hoagland nutrient solution was used during the process (Hoagland 1950). When the third leaves of maize plants were fully expanded, hydroponics models were constructed using three groups: group A (Hoagland), group B (Hoagland + PEG), group C (Hoagland + PEG + 4 mg/kg S-ABA). Samples were taken at 2, 4, 8, 12, and 24 h after treatment; they were washed with double-distilled water and then frozen in liquid nitrogen for 2 min and vacuum freeze-dried before being stored at – 80 °C until analysis.

### Extraction and Quantification of Leaf ABA

The method for the extraction of leaf ABA was adapted from the study of Wang et al. (1998). Five hundred milligrams of leaf tissue were macerated in liquid nitrogen and mixed with 5 ml 80% (v/v) methanol containing 0.1% butylated hydroxytoluene (BHT) as the antioxidant. The extract was incubated at 4 °C for 4 h and subsequently centrifuged for 15 min at 12,000 $\times$ g at the same temperature. The supernatant fluid was collected, and an immunoassay was carried out according to the ELISA kit procedure (the ABA Enzyme Immunoassay Test Kit was provided by China Agricultural University).

### Determination of Proline Content

Free proline content was measured using ninhydrin colorimetry (Abraham et al. 2010). Fresh maize leaf material (0.5 g) from each replicate within each treatment was triturated in 5 ml of 3% sulfosalicylic acid solution. Two milliliters of the filtrate was reacted with 3 ml acid ninhydrin solution and 3 ml of glacial acetic acid in a test tube; the sample mixture was then incubated at 100 °C for 60 min. After cooling, 4 ml of toluene was added to the mixture and it was mixed

vigorously for 1–2 min. The absorbance of the colored portion containing toluene was read at 520 nm using toluene as the blank.

### Estimation of Soluble Sugar Content

Leaf total soluble sugar concentrations were evaluated at 620 nm using a spectrophotometer according to the anthrone method (Fales 1951). Fresh leaves (0.5 g) were

put into 15 ml of distilled water and boiled in a water bath for 20 min. After cooling, 5 ml of anthrone was added to 0.1 ml of boiled sample. Then, 3 ml of boiled sample was transferred to a cuvette and the absorbance was read. Finally, the amount of total soluble sugar in the samples was quantified using a standard glucose curve.

### Determination of Hydrogen Peroxide Content

Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) content in leaves was determined according to the method of Prasad et al. (1994). Samples (0.5 g) were homogenized in an ice bath with 2 ml of 0.1% (w/v) trichloroacetic acid. The homogenate was centrifuged at 10,000 $\times$ g for 10 min, and 1 ml of the supernatant was added to 1 ml of 10 mM potassium phosphate buffer (pH 7.0) and 2 ml of 1 M KI. The absorbance of the supernatant was measured at 390 nm.

### Estimation of Malondialdehyde Content

Malondialdehyde (MDA) content of the maize leaves was analyzed using the thiobarbituric acid (TBA) method (Cakmak and Horst 2006). Fresh leaf samples (0.2 g) were triturated in a 10 ml solution of 0.1% TCA. The sample was then centrifuged at 12,000 $\times$ g for 10 min, and the supernatant was separated. To 1 ml of the supernatant, 4 ml of 20% TCA solution was added, which contained 0.5% TBA solution. Subsequently, the sample was heated in a water bath at 95 °C for 40 min and then cooled immediately on ice for 5 min. The cooled samples were centrifuged for 15 min at 3000 $\times$ g at 4 °C, and then the absorbance of the supernatant was read at 532 nm, 600 nm, and 450 nm.

### Enzyme Extraction and Activity

Enzyme extraction was carried out by immersing 0.5 g of leaves in liquid nitrogen, to which was added 10 ml of

extraction buffer made up of 0.05 M potassium phosphate buffer (pH 7.8), 0.1 mM EDTA, 0.3% Triton-X100, and 4% polyvinylpyrrolidone (PVP); this was repeated three times. The extract was centrifuged at  $15,000\times g$  for 20 min at 4 °C, and the supernatant was collected and stored at – 80 °C during the analysis period. The collected supernatants were used in all enzyme analyses. The activity of enzymes was expressed in milligrams (mg) of proteins, which were determined by the Bradford (1976) method, using the standard curve of bovine serum albumin (BSA).

The activity of superoxide dismutase (SOD, EC 1.15.1.1) was evaluated by its ability to inhibit the photoreduction of nitroblue tetrazolium (NBT), using the method of Giannopolitis and Ries (1977). Measurements were taken at 560 nm and one unit of SOD corresponded to the amount of enzyme capable of inhibiting 50% of NBT photoreduction under experimental conditions. Catalase activity (CAT, EC 1.11.1.6) was determined by  $H_2O_2$  consumption at 240 nm during 3 min (Havir and McHale 1987). Peroxidase (POD, EC 1.11.1.7) activity was determined by the guaiacol method (Lin and Wang 2002). The absorbance of the reaction was read at 420 nm with a 30 s interval up to 2 min; this used the absorbance change 0.01 as a POD activity. Ascorbate peroxidase activity (APX EC 1.11.1.11) was assayed according to Nakano and Asada (1981). The change in absorbance after adding  $H_2O_2$  was read at 290 nm for 2 min at 30 s intervals. Polyphenol oxidase activity (PPO, EC 1.10.3.2) was assessed through the method proposed by Mohammadi and Kazemi (2002) with some modifications. The activity of phenylalanine ammonia lyase (PAL, EC 4.3.1.5) was determined by the formation of cinnamic acid at 290 nm (Cahill and McComb 1992).

## RT-qPCR

Total RNA was extracted using the Ominiplant Plant Kit (DNase I) (CWBIO) from samples homogenized in liquid nitrogen using a pestle and mortar (2  $\times$  3 repetitions). cDNA was synthesized using a HiFiScript cDNA Synthesis Kit (CWBIO). cDNA (20 $\times$  diluted) was mixed with the UltraSYBR Mixture (Low ROX; CWBIO) and 200 nM of respective primers to a final volume of 25  $\mu$ l. The RT-qPCR was performed using a LightCycler® 96 (Roche Applied Science). The qPCR program was set to an initial denaturation (10 min, 95 °C), followed by 40 cycles of primer denaturation (15 s, 95 °C), annealing, and elongation (60 s, 60 °C). The relative content of RNA was calculated according to the method of Hellemans et al. (2007). *EF1 $\alpha$*  was used as the reference gene. Primers were designed according to sequences retrieved from the NCBI database (Lamesch et al. 2012) using the Primer3 program (Untergasser et al. 2007). Primer sequences are shown in Table S1.

## Statistical Analysis

Data were analyzed using the Microsoft Excel 2010 and SPSS 20.0 statistical packages. Origin 2018 was used for graphical presentation of the data. A factorial experiment based on randomized complete block design was carried out with three replicates ( $n = 3$ ). Duncan's multiple range test ( $p < 0.05$ ) was used to compare the means.

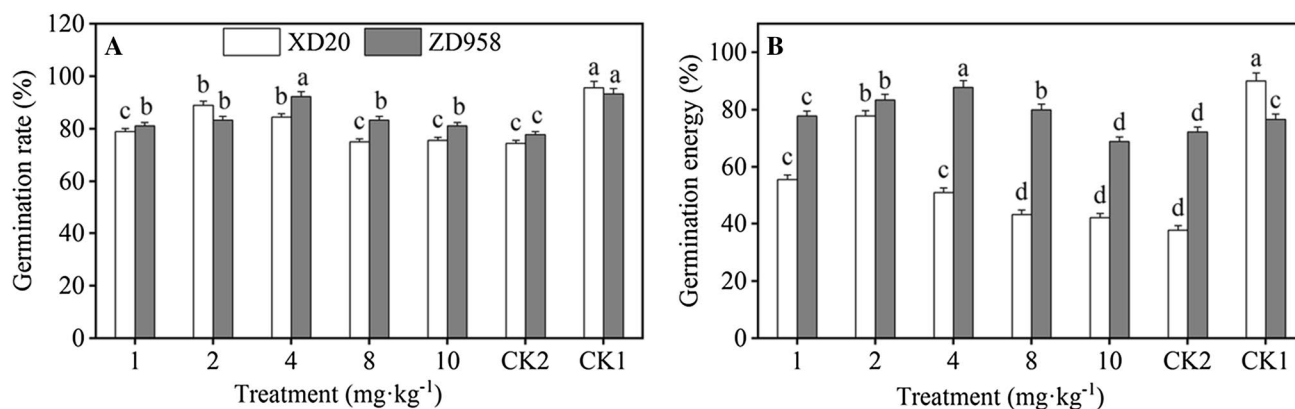
## Results

### Effects of S-ABA Seed Soaking on the Biological Character of Maize Seeds Under Drought Stress

Compared with CK1, drought stress significantly inhibited germination in the two maize varieties. The GR and GE of XD20 and ZD958 decreased by 23.19%, 18.81%, 41.96%, and 22.09%, respectively. Under drought conditions, seed soaking with different concentrations of S-ABA could increase the GR and GE of seeds; furthermore, with an increase in soaking concentration, the trend first increased then decreased (Fig. 1). For GR, all treatments (except the 10 mg/kg treatment) of ZD958 reached significant levels compared to CK2, whereas XD20 reached significant levels only at concentrations of 1, 2 and 4 mg/kg ( $p < 0.05$ ). These results suggest that soaking seeds with suitable concentrations of S-ABA can effectively mitigate the effect of drought stress on the germination of maize seeds, with the concentration of 4 mg/kg being optimum ( $p < 0.05$ ; Fig. 1).

### Effects of S-ABA Treatment on the Growth of Maize Seedlings Under Drought Stress

Regarding water stress, root and bud growth of both varieties was drastically reduced; this was expressed as a sharp decrease in root and shoot fresh and dry weight, although the drought stress on ZD958 was less evident ( $p < 0.05$ ). Under drought stress, the root length, bud length, lateral root number, root fresh weight, bud fresh weight, root dry weight, and bud dry weight of the two maize varieties when soaked with S-ABA were significantly increased ( $p < 0.05$ ). Analysis of variance results showed that soaking the seeds in different concentrations of S-ABA mitigated the effect of drought stress on maize seedlings to varying degrees. When compared with the drought control, the treatment of S-ABA at the concentration of 4 mg/kg was most significant ( $p < 0.05$ ) (Table 1). Meanwhile, the results showed that 4 mg/kg of S-ABA was most effective for alleviating drought ( $p < 0.05$ ). Compared to CK2, the average root length, bud length, lateral root number, root fresh weight, bud fresh weight, root dry weight, and bud dry weight of ZD958 increased 48.97%, 82.41%, 38.62%, 44.14%, 73.96%, 16.18%, and 73.68%,



**Fig. 1** Effects of S-ABA seed soaking on **a** germination rate and **b** germination energy of maize seeds under drought stress. Means followed by the same letters for the treatments do not differ according to Duncan's test at 5% significance ( $p < 0.05$ ). Capped bars above means

represent  $\pm$  SE of three replicates. Small alphabetical letters above means denote the significant differences among treatment within two maize hybrids at  $p < 0.05$ . CK1, control (irrigate); CK2, control (drought)

**Table 1** Effects of S-ABA treatment on the growth of maize seedlings under drought stress

Variety	Concentration (mg/kg)	Root length (cm)	Bud length (cm)	Lateral root number	Root fresh weight (g)	Bud fresh weight (g)	Root dry weight (g)	Bud dry weight (g)
ZD 958	1	6.82 $\pm$ 0.14 <sup>d</sup>	3.19 $\pm$ 0.34 <sup>d</sup>	3.50 $\pm$ 0.30 <sup>b,c</sup>	1.64 $\pm$ 0.01 <sup>d,e</sup>	1.25 $\pm$ 0.26 <sup>b,c</sup>	0.74 $\pm$ 0.10 <sup>b</sup>	0.26 $\pm$ 0.05 <sup>b,c,d</sup>
	2	7.15 $\pm$ 0.31 <sup>c,d</sup>	3.66 $\pm$ 0.36 <sup>c,d</sup>	3.85 $\pm$ 0.25 <sup>b,c</sup>	1.88 $\pm$ 0.08 <sup>b,c</sup>	1.41 $\pm$ 0.27 <sup>b,c</sup>	0.78 $\pm$ 0.07 <sup>b</sup>	0.30 $\pm$ 0.04 <sup>a,b,c</sup>
	4	8.03 $\pm$ 1.20 <sup>b</sup>	4.98 $\pm$ 0.13 <sup>b</sup>	4.81 $\pm$ 0.59 <sup>a</sup>	2.09 $\pm$ 0.14 <sup>b</sup>	1.67 $\pm$ 0.29 <sup>b</sup>	0.79 $\pm$ 0.06 <sup>b</sup>	0.33 $\pm$ 0.01 <sup>a,b</sup>
	8	7.47 $\pm$ 0.34 <sup>c</sup>	3.84 $\pm$ 0.19 <sup>c</sup>	4.17 $\pm$ 0.31 <sup>b</sup>	1.93 $\pm$ 0.10 <sup>b,c</sup>	1.29 $\pm$ 0.43 <sup>b,c</sup>	0.75 $\pm$ 0.08 <sup>b</sup>	0.27 $\pm$ 0.09 <sup>b,c,d</sup>
	10	7.13 $\pm$ 1.07 <sup>c,d</sup>	3.56 $\pm$ 0.34 <sup>c,d</sup>	3.73 $\pm$ 0.32 <sup>b,c</sup>	1.82 $\pm$ 0.06 <sup>c,d</sup>	1.15 $\pm$ 0.20 <sup>b,c</sup>	0.74 $\pm$ 0.10 <sup>b</sup>	0.22 $\pm$ 0.03 <sup>c,d</sup>
	CK2	5.39 $\pm$ 1.11 <sup>e</sup>	2.73 $\pm$ 0.28 <sup>e</sup>	3.47 $\pm$ 0.24 <sup>e</sup>	1.45 $\pm$ 0.14 <sup>e</sup>	0.96 $\pm$ 0.15 <sup>c</sup>	0.68 $\pm$ 0.02 <sup>c</sup>	0.19 $\pm$ 0.04 <sup>d</sup>
	CK1	9.16 $\pm$ 0.48 <sup>a</sup>	6.68 $\pm$ 0.31 <sup>a</sup>	3.20 $\pm$ 0.46 <sup>e</sup>	2.89 $\pm$ 0.24 <sup>a</sup>	2.74 $\pm$ 0.29 <sup>a</sup>	0.83 $\pm$ 0.07 <sup>a</sup>	0.37 $\pm$ 0.05 <sup>a</sup>
XD 20	1	6.22 $\pm$ 1.36 <sup>d</sup>	2.78 $\pm$ 0.22 <sup>d</sup>	3.64 $\pm$ 0.39 <sup>b,c,d</sup>	1.57 $\pm$ 0.22 <sup>b,c</sup>	1.06 $\pm$ 0.09 <sup>c</sup>	0.72 $\pm$ 0.03 <sup>b,c</sup>	0.24 $\pm$ 0.04 <sup>b</sup>
	2	6.56 $\pm$ 0.48 <sup>c</sup>	3.14 $\pm$ 0.14 <sup>c,d</sup>	3.83 $\pm$ 0.25 <sup>b,c</sup>	1.71 $\pm$ 0.14 <sup>b</sup>	1.22 $\pm$ 0.12 <sup>b</sup>	0.74 $\pm$ 0.04 <sup>b</sup>	0.25 $\pm$ 0.01 <sup>b</sup>
	4	7.12 $\pm$ 1.26 <sup>b</sup>	4.05 $\pm$ 0.31 <sup>b</sup>	4.46 $\pm$ 0.10 <sup>a</sup>	1.42 $\pm$ 0.02 <sup>b,c</sup>	0.95 $\pm$ 0.01 <sup>c,d</sup>	0.72 $\pm$ 0.01 <sup>b,c</sup>	0.23 $\pm$ 0.02 <sup>b,c</sup>
	8	6.64 $\pm$ 0.65 <sup>c</sup>	3.58 $\pm$ 0.57 <sup>b,c</sup>	3.95 $\pm$ 0.26 <sup>b</sup>	1.38 $\pm$ 0.01 <sup>b,c</sup>	0.88 $\pm$ 0.02 <sup>d,e</sup>	0.69 $\pm$ 0.02 <sup>b,c</sup>	0.21 $\pm$ 0.01 <sup>b,c</sup>
	10	6.16 $\pm$ 1.41 <sup>d</sup>	3.27 $\pm$ 0.24 <sup>c,d</sup>	3.64 $\pm$ 0.39 <sup>b,c,d</sup>	1.29 $\pm$ 0.17 <sup>c</sup>	0.86 $\pm$ 0.07 <sup>d,e</sup>	0.70 $\pm$ 0.03 <sup>b,c</sup>	0.19 $\pm$ 0.03 <sup>c,d</sup>
	CK2	5.25 $\pm$ 1.38 <sup>e</sup>	2.28 $\pm$ 0.11 <sup>e</sup>	3.40 $\pm$ 0.13 <sup>c,d</sup>	1.27 $\pm$ 0.34 <sup>c</sup>	0.77 $\pm$ 0.09 <sup>e</sup>	0.63 $\pm$ 0.11 <sup>c</sup>	0.17 $\pm$ 0.01 <sup>d</sup>
	CK1	10.31 $\pm$ 0.75 <sup>a</sup>	7.26 $\pm$ 0.25 <sup>a</sup>	3.16 $\pm$ 0.10 <sup>d</sup>	2.92 $\pm$ 0.24 <sup>a</sup>	2.76 $\pm$ 0.11 <sup>a</sup>	0.85 $\pm$ 0.06 <sup>a</sup>	0.39 $\pm$ 0.08 <sup>a</sup>

Means followed by the same letters for the treatments do not differ according to Duncan's test at 5% significance ( $p < 0.05$ ). Each value indicates the treatment average  $\pm$  SE ( $n = 3$ )

CK1 control (irrigate), CK2 control (drought), ZD958 'Zhengdan 958' (drought tolerant), XD20 'Xundan 20' (drought sensitive)

respectively; whereas those of XD20 increased 35.62%, 77.63%, 31.18%, 11.81%, 23.37%, 14.28%, and 35.29%, respectively. Moreover, the results from ZD958 were markedly superior to those of XD20 ( $p < 0.05$ ).

### Effects of S-ABA Treatment on Endogenous ABA Content in Maize Leaves Under Drought Stress

In all treatments, analysis of variance showed that there was essentially no difference in endogenous ABA content between the two cultivars under the normal watering treatment, whereas endogenous ABA content in the leaves of the two maize

cultivars treated with PEG and PEG + ABA presented a sharp rise ( $p < 0.05$ ). After treatment for 8 h, the ABA content of endogenous hormone in ZD958 reached its maximum, which was 84.59% higher than that in the drought control. The ABA content of endogenous hormone in XD20 reached a maximum after treatment for 24 h, which was an increase of 75.80% (Table 2).



**Table 2** Effects of S-ABA treatment on endogenous abscisic acid (ABA) content in maize leaves under drought stress

Treatment (h)	2	4	8	12	24
Endogenous abscisic acid (ABA) content (ng mL <sup>-1</sup> FW)					
XD20 + PEG	146.68 ± 11.00 <sup>c</sup>	100.35 ± 10.03 <sup>c</sup>	125.38 ± 5.18 <sup>c</sup>	109.69 ± 1.94 <sup>c</sup>	141.07 ± 7.77 <sup>c</sup>
ZD958 + PEG	169.09 ± 10.05 <sup>b,c</sup>	188.89 ± 6.38 <sup>b</sup>	129.86 ± 9.06 <sup>c</sup>	117.16 ± 3.56 <sup>c</sup>	147.80 ± 7.12 <sup>c</sup>
XD20 + PEG + ABA	176.19 ± 10.68 <sup>b</sup>	172.45 ± 12.63 <sup>b</sup>	190.39 ± 6.47 <sup>b</sup>	172.83 ± 3.58 <sup>b</sup>	200.48 ± 4.53 <sup>a</sup>
ZD958 + PEG + ABA	222.89 ± 5.18 <sup>a</sup>	236.34 ± 10.35 <sup>a</sup>	239.71 ± 1.94 <sup>a</sup>	217.29 ± 8.41 <sup>a</sup>	178.81 ± 4.21 <sup>b</sup>
XD20	77.56 ± 1.98 <sup>d</sup>	76.06 ± 2.33 <sup>d</sup>	76.44 ± 1.98 <sup>d</sup>	75.32 ± 2.70 <sup>d</sup>	74.19 ± 0.99 <sup>d</sup>
ZD958	78.68 ± 2.62 <sup>d</sup>	80.17 ± 1.87 <sup>d</sup>	77.56 ± 0.99 <sup>d</sup>	81.29 ± 4.54 <sup>d</sup>	80.55 ± 2.33 <sup>d</sup>

Each value indicates the treatment average ± SE ( $n=3$ ). One-way analyses of variance (ANOVA) were employed to test the effects of drought stress, S-ABA and their interaction, and means were separated using Duncan's multiple range tests. Differences were considered significant at  $p < 0.05$ . ZD958 'Zhengdan958' (drought tolerant), XD20 'Xundan 20' (drought sensitive); PEG, 15% w/v, MW 6000; S-ABA, 4 mg kg<sup>-1</sup> S-ABA

### Effects of S-ABA Treatment on Hydrogen Peroxide (H<sub>2</sub>O<sub>2</sub>) Content in Maize Leaves Under Drought Stress

Compared with the control, the treatment with PEG and PEG + ABA significantly increased the H<sub>2</sub>O<sub>2</sub> content; the PEG + ABA treatment showed the highest H<sub>2</sub>O<sub>2</sub> content, and the treatment of XD20 showed greater sensitivity ( $p < 0.05$ ). After 12 h of stress, XD20 + PEG + ABA showed the highest content of H<sub>2</sub>O<sub>2</sub> (Table 3). The ZD958 + PEG + ABA exhibited significantly higher H<sub>2</sub>O<sub>2</sub> content at 8 h under water stress, followed by XD20 + PEG, and ZD958 + PEG. There was no difference among the irrigation treatments ( $p < 0.05$ ).

### Effects of S-ABA Treatment on MDA Content of Maize Leaves Under Drought Stress

Malondialdehyde level in maize leaves showed a significant tendency to increase in both the PEG and PEG + ABA treatments compared with the control group ( $p < 0.05$ ). However, MDA in XD20 leaves increased faster than that in ZD958 leaves ( $p < 0.05$ ). The MDA of maize in the PEG and PEG + ABA groups was higher than that in the control

group in the 2 h treatment, but the difference was insignificant ( $p < 0.05$ ). From 4 to 24 h, there was a significant difference (elevation) in MDA content between maize leaves treated with drought and normal growth; however, the rate of increment was higher in XD20 at 12 h ( $p < 0.05$ ). Results also showed that MDA content reached a maximum at 24 h. After ABA treatment, the decrease in MDA content in leaves of the two maize cultivars also reached a significant level after 12 h and 24 h of stress ( $p < 0.05$ ; Table 4).

### Effects of S-ABA Treatment on Osmotic Substance Content in Maize Leaves Under Drought Stress

As a result, the proline content showed an increasing tendency with stress time and the difference was significant in each time period ( $p < 0.05$ ). After treatment with S-ABA for 4 h, the difference in treatment between PEG + ABA and PEG was significant. The proline content of ZD958 was higher than that of XD20 in all treatments ( $p < 0.05$ ; Fig. 2a). After treatment for 24 h, the content of proline in the PEG + ABA treatment reached a maximum; ZD 958 increased by 28.23% compared with the PEG treatment, and XD20 increased by 27.41%.

**Table 3** Effects of S-ABA treatment on hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) content in maize leaves under drought stress

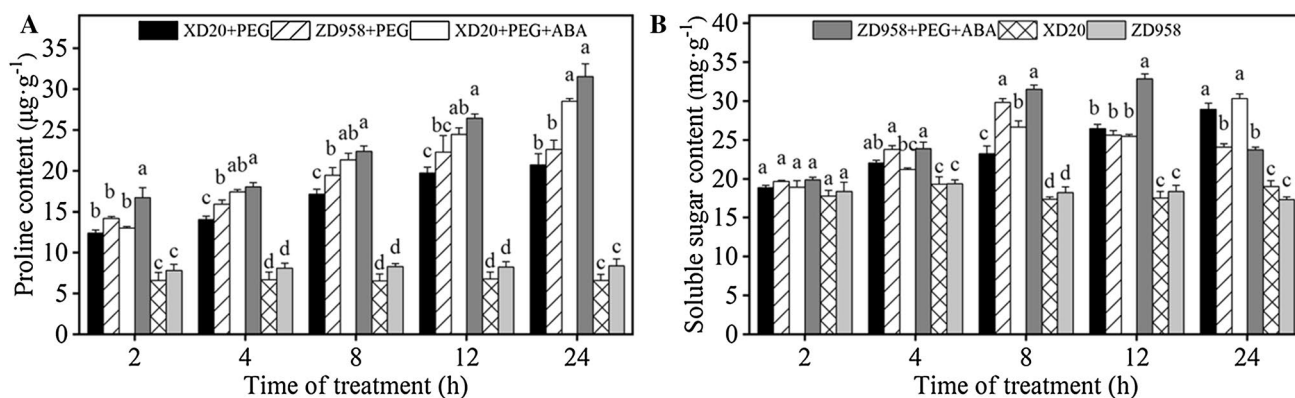
Treatment (h)	2	4	8	12	24
Hydrogen peroxide (H <sub>2</sub> O <sub>2</sub> ) content (μmol g <sup>-1</sup> FW)					
XD20 + PEG	30.41 ± 2.25 <sup>c</sup>	33.47 ± 1.57 <sup>c</sup>	41.09 ± 2.29 <sup>d</sup>	54.98 ± 2.27 <sup>b</sup>	51.48 ± 1.22 <sup>b</sup>
ZD958 + PEG	25.42 ± 0.92 <sup>d</sup>	31.21 ± 1.34 <sup>c</sup>	49.13 ± 1.19 <sup>c</sup>	40.25 ± 1.44 <sup>d</sup>	38.17 ± 2.60 <sup>c</sup>
XD20 + PEG + ABA	43.71 ± 1.11 <sup>a</sup>	53.23 ± 2.49 <sup>a</sup>	61.24 ± 2.06 <sup>b</sup>	71.69 ± 1.34 <sup>a</sup>	64.31 ± 2.79 <sup>a</sup>
ZD958 + PEG + ABA	33.82 ± 1.57 <sup>b</sup>	38.78 ± 1.93 <sup>b</sup>	69.01 ± 1.20 <sup>a</sup>	51.18 ± 1.78 <sup>c</sup>	45.61 ± 1.42 <sup>b</sup>
XD20	18.46 ± 0.88 <sup>e</sup>	20.38 ± 1.05 <sup>d</sup>	20.16 ± 1.93 <sup>e</sup>	19.11 ± 2.03 <sup>e</sup>	19.79 ± 1.48 <sup>d</sup>
ZD958	17.85 ± 1.61 <sup>e</sup>	17.56 ± 1.26 <sup>d</sup>	18.82 ± 1.58 <sup>e</sup>	18.76 ± 2.03 <sup>e</sup>	19.39 ± 0.78 <sup>d</sup>

Each value indicates the treatment average ± SE ( $n=3$ ). One-way analyses of variance (ANOVA) were employed to test the effects of drought stress, S-ABA and their interaction, and means were separated using Duncan's multiple range tests. Differences were considered significant at  $p < 0.05$ . ZD958 'Zhengdan958' (drought tolerant), XD20 'Xundan 20' (drought sensitive); PEG, 15% w/v, MW 6000; S-ABA, 4 mg kg<sup>-1</sup> S-ABA

**Table 4** Effects of S-ABA treatment on malondialdehyde (MDA) content of maize leaves under drought stress

Treatment (4)	2	4	8	12	24
Malondialdehyde (MDA) content (nmol g <sup>-1</sup> FW)					
XD20 + PEG	13.92 ± 0.34 <sup>a</sup>	18.49 ± 1.63 <sup>a</sup>	22.84 ± 0.73 <sup>a</sup>	28.09 ± 2.62 <sup>a</sup>	32.58 ± 1.24 <sup>a</sup>
ZD958 + PEG	15.04 ± 1.86 <sup>a</sup>	18.5 ± 1.59 <sup>a</sup>	20.59 ± 0.74 <sup>a,b</sup>	23.21 ± 1.77 <sup>b</sup>	27.20 ± 1.32 <sup>b</sup>
XD20 + PEG + ABA	12.31 ± 1.25 <sup>a</sup>	14.94 ± 0.92 <sup>a,b</sup>	19.15 ± 0.45 <sup>a,b</sup>	22.05 ± 1.34 <sup>b</sup>	24.19 ± 2.24 <sup>b,c</sup>
ZD958 + PEG + ABA	13.22 ± 0.67 <sup>a</sup>	15.9 ± 1.60 <sup>a</sup>	18.24 ± 0.27 <sup>b</sup>	20.37 ± 2.38 <sup>c</sup>	23.20 ± 0.50 <sup>c</sup>
XD20	10.55 ± 2.86 <sup>a</sup>	9.97 ± 1.04 <sup>c</sup>	11.85 ± 2.06 <sup>c</sup>	11.39 ± 0.66 <sup>d</sup>	11.76 ± 0.92 <sup>d</sup>
ZD958	10.17 ± 1.33 <sup>a</sup>	10.88 ± 1.10 <sup>b,c</sup>	11.93 ± 1.83 <sup>c</sup>	12.38 ± 1.34 <sup>d</sup>	12.07 ± 1.02 <sup>d</sup>

Each value indicates the treatment average ± SE ( $n=3$ ). One-way analyses of variance (ANOVA) were employed to test the effects of drought stress, S-ABA and their interaction, and means were separated using Duncan's multiple range tests. Differences were considered significant at  $p < 0.05$ . ZD958 'Zhengdan958' (drought tolerant), XD20 'Xundan 20' (drought sensitive); PEG, 15% w/v, MW 6000; S-ABA, 4 mg kg<sup>-1</sup> S-ABA



**Fig. 2** Effects of S-ABA treatment on **a** proline and **b** soluble sugar content in maize leaves under drought stress. Means followed by the same letters for the treatments do not differ according to Duncan's test at 5% significance ( $p < 0.05$ ). Capped bars above means repre-

sent ± SE of three replicates. Small alphabetical letters above means denote the significant differences among treatment with in two maize hybrids at  $p < 0.05$

Analysis of variance showed that drought stress increased the soluble sugar content of maize leaves, and the application of exogenous ABA further increased the soluble sugar content. There were no significant differences between all experiments after 2 h of treatment ( $p < 0.05$ ). After 8 h of stress, the soluble sugar content of XD20 + PEG + ABA treatment peaked. After 12 h of stress, ZD958 + PEG + ABA exhibited the highest soluble sugar contents followed by the treatments XD20 + PEG, ZD958 + PEG, and XD958 + PEG + ABA. With increasing time (24 h), there was an inversion in the results, with XD20 + PEG + ABA showing a significantly higher content, followed by XD20 + PEG (Fig. 2b).

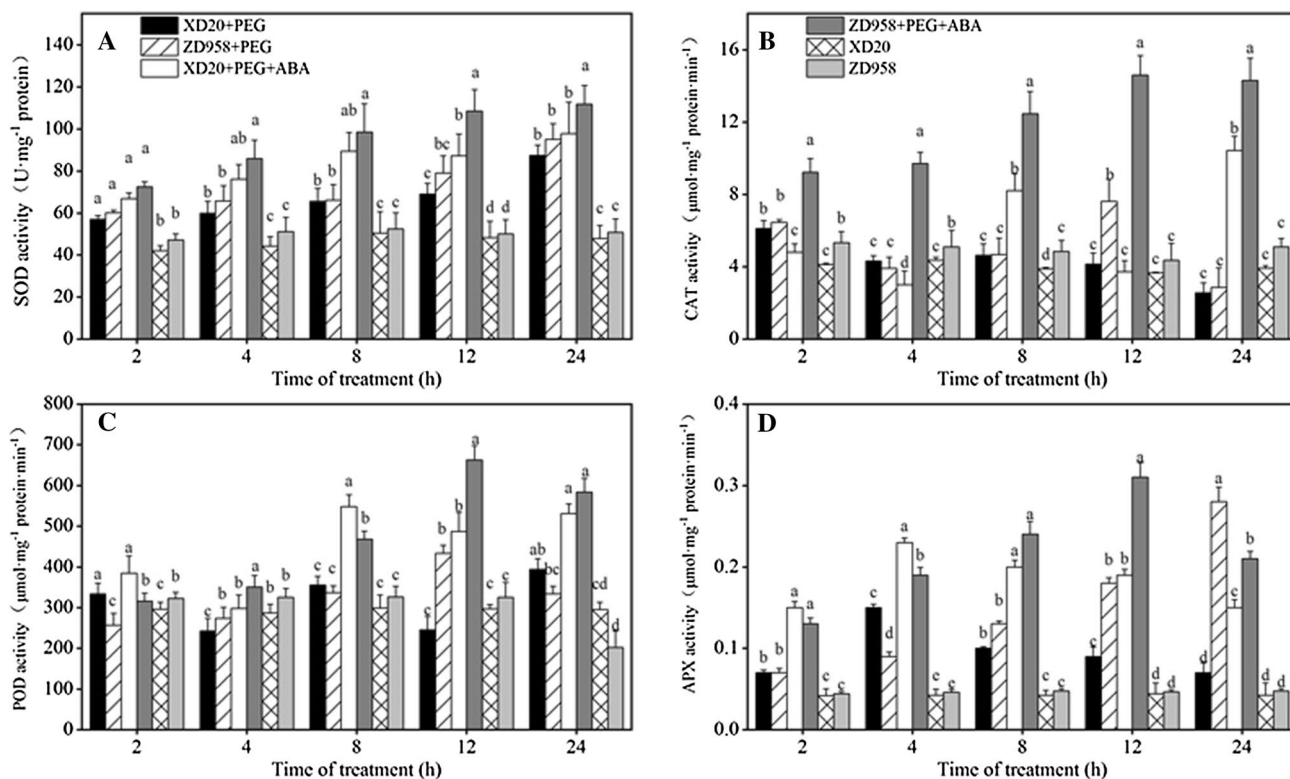
### Effects of S-ABA Treatment on Enzyme Activity in Maize Leaves Under Drought Stress

The increase in drought stress time resulted in higher enzyme activity in both ZD958 and XD20. However, with normal watering there was no significant change ( $p < 0.05$ ). With increasing stress time, the activity of protective enzymes

(including SOD, CAT, POD, APX, PPO, and PAL) in the leaves initially rose and then declined (Figs. 3, 4).

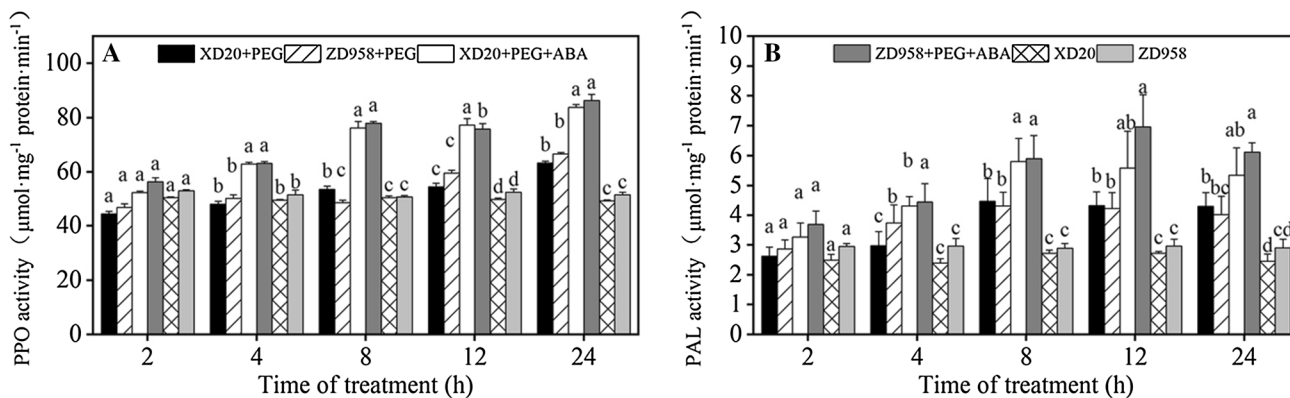
No significant differences were observed in the analysis of SOD activity among treatments during the first 2 h of stress (Fig. 3a). After the fourth hour of stress, there was a significant increase in SOD activity in the PEG and PEG + ABA treatments of ZD958 ( $p < 0.05$ ). Furthermore, the activity of SOD peaked at 12 h, and was 19.71% higher than that of the PEG control. However, XD20 showed a significant difference after 12 h of treatment; SOD activity peaked after 8 h of treatment, which was a 12.74% increase compared with the PEG control ( $p < 0.05$ ; Fig. 3a).

Under 2 h of drought stress, S-ABA application led to an increase in CAT activity in the leaves of ZD958 under water stress conditions; CAT activity was higher in ZD958 than in XD20 ( $p < 0.05$ ). The CAT activity of ZD958 gradually increased and peaked at 12 h, which was an increase of 91.47% over the drought control ( $p < 0.05$ ). However, the CAT activity of XD20 + PEG + ABA was lower than that of XD20 + PEG in the first 4 h stress treatment. XD20



**Fig. 3** Effects of S-ABA treatment on the activities of **a** superoxide dismutase (SOD), **b** catalase (CAT), **c** peroxidase (POD) and **d** ascorbate peroxidase (APX) maize leaves under drought stress. Means followed by the same letters for the treatments do not differ according to

Duncan’s test at 5% significance ( $p < 0.05$ ). Capped bars above means represent  $\pm$ SE of three replicates. Small alphabetical letters above means denote the significant differences among treatment with in two maize hybrids at  $p < 0.05$



**Fig. 4** Effects of S-ABA treatment on the activities of **a** polyphenol oxidase (PPO) and **b** phenylalanine ammonia lyase (PAL) maize leaves under drought stress. Means followed by the same letters for the treatments do not differ according to Duncan’s test at 5% signifi-

cance ( $p < 0.05$ ). Capped bars above means represent  $\pm$ SE of three replicates. Small alphabetical letters above means denote the significant differences among treatment with in two maize hybrids at  $p < 0.05$

showed a significant difference only after 8 h and 24 h. CAT activity reached a maximum at 24 h of stress, which was an increase of 307.42% compared with the XD20 + PEG treatment. After 24 h of stress, ZD958 + PEG + ABA increased by 401.40% compared with ZD958 + PEG. Furthermore, the

CAT activity of XD20 initially increased and then decreased and finally increased again after treatment with S-ABA ( $p < 0.05$ ; Fig. 3b).

In the evaluation of POD enzyme activity, the treatments showed no obvious differences at 4 h of stress



(Fig. 3c). At 12 h in the stressed treatment, ZD958 showed superior POD activity with S-ABA application and an increase of 52.89% compared to the PEG + ZD958 treatment ( $p < 0.05$ ). After 24 h of stress, S-ABA application also elevated the POD enzyme activity of XD20 + PEG + ABA to values similar to those of ZD958 + PEG + ABA. In addition, the treatment XD20 + PEG + ABA showed a POD activity increase of 35.02% compared with XD20 + PEG (Fig. 3c).

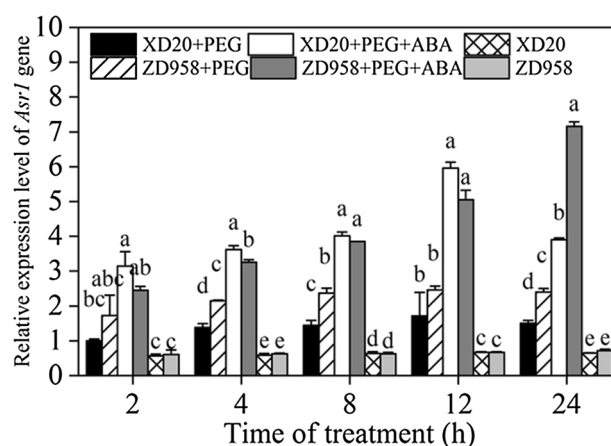
After 2 h of stress, the treatments PEG and PEG + ABA of the two maize cultivars had greater activity of APX enzyme than under the remaining treatments. In addition, the PEG + ABA treatment differed significantly ( $p < 0.05$ ). ZD958 had the highest activity of APX enzyme after treatment for 12 h, which was 72.22% higher than that of the drought treatment. These figures for XD20 peaked after treatment for 4 h, an increase of 53.33%.

After stress treatment for 12 h, the stressing treatments (PEG and PEG with ABA application) for both hybrids produced significantly more pronounced PPO activity than the irrigated treatments ( $p < 0.05$ ). After 2 h of stress treatment, the difference between the treatments was not significant. After 4 h of stress treatment, the application of S-ABA significantly increased the PPO enzyme activity of the two maize varieties. After 24 h of stress, XD20 + PEG + ABA and ZD958 + PEG + ABA showed greatly elevated means; compared with the PEG treatment, they increased by 32.27% and 29.41%, respectively (Fig. 4a).

Regarding PAL means activity, there were no significant differences among the treatments given 2 h of water stress ( $p < 0.05$ ). After 4 h of stress treatment, the application of S-ABA showed significant differences. ZD958 + PEG + ABA had the highest activity of PAL enzyme at 12 h, which was 64.54% higher than that in the PEG treatment ( $p < 0.05$ ). However, XD20 + PEG + ABA had the highest activity of PAL enzyme at 8 h, which was 29.98% higher than that in the PEG treatment ( $p < 0.05$ ; Fig. 4b).

### Effects of S-ABA Treatment on *Asr1* Gene-Expressive Level in Maize Leaves Under Drought Stress

Water stress increased the gene expressive level of *Asr1* in both cultivars. Compared with drought stress, the expression levels of *Asr1* in the two varieties after exogenous application of S-ABA were up-regulated at different time periods ( $p < 0.05$ ). Among them, the gene expressive level of XD20 *Asr1* was the highest after the 12-h treatment with S-ABA, whereas that of ZD958 reached a maximum at 24 h ( $p < 0.05$ ). In contrast, the increase of *Asr1* expression level in ZD958 was greater than that in XD20 (Fig. 5).



**Fig. 5** Effects of S-ABA treatment on *Asr1* gene-expressive quantity in maize leaves under drought stress. Means followed by the same letters for the treatments do not differ according to Duncan's test at 5% significance ( $p < 0.05$ ). Capped bars above means represent  $\pm$  SE of three replicates. Small alphabetical letters above means denote the significant differences among treatment with in two maize hybrids at  $p < 0.05$

## Discussion

Water stress is one of the most serious environmental factors limiting plant growth and development. The relationship between ABA accumulation and stress resistance under stress conditions has become an important area of stress research. Our results indicated that the GE, GR, and seedling biomass of the two maize varieties decreased significantly under drought stress ( $p < 0.05$ ). This observation agrees well with the results of Yongsheng et al. (2016). Abscisic acid is an important phytohormone, not only because of its regulatory functions in physiological and developmental processes but also because it is the main regulator in adaptive responses to drought stress (Jiang and Zhang 2002; Lou et al. 2017). The results showed that the soaking of seedlings with S-ABA significantly promoted maize seed germination under drought stress, but there was a significant inhibitory effect at high concentrations. Among the treatments tested, 4 mg/kg S-ABA had the best effect ( $p < 0.05$ ; Fig. 1). These results suggest that the expression of maize drought resistance-related genes can be activated by 4 mg/kg of S-ABA, thereby alleviating the inhibition of drought on maize seed germination and seedling biomass accumulation, and increasing the germination rate of maize seeds under drought stress. This is consistent with the results reported by Gang et al. (2017), who reported that S-ABA spray can alleviate the inhibition of drought on the growth of maize seedlings. However, the different maize varieties had different responses to drought stress with XD20 being more sensitive than ZD958 (Table 1).

ABA is a major regulator of drought stress, and some studies report an increase in endogenous ABA in drought-tolerant maize under drought stress (De Souza et al. 2014). ABA balance between the roots and leaves has also been reported to affect plant moisture status. Water stress increases plant-endogenous ABA content, and the external application of ABA further promotes the synthesis of endogenous ABA to increase total levels; this has been confirmed in wheat, sugarcane, and other crops (Li et al. 2017; Ma et al. 2016; Virilouvet et al. 2011). Our results showed that endogenous ABA content increased significantly with the prolongation of water stress ( $p < 0.05$ ). Similarly, Alvarez et al. (2014) report that in wheat, lower ABA accumulation can be found in drought-tolerant genotypes towards the end of the stress treatment. Compared with the drought treatment, exogenous application of S-ABA increased ABA content, in agreement with the results of Kizis and Pagès (2002) (Table 2). Furthermore, Zhang and Xie (2002) show that ABA positively regulates the content of endogenous hormones; this may be an important reason why ABA can improve drought resistance in maize; however, further research is needed to confirm this finding.

A noticeable fact concerning hydrogen peroxide ( $H_2O_2$ ) was that, with the increase of drought stress time, PEG and PEG + ABA treatments showed higher  $H_2O_2$  content; with ZD958 under this same treatment, a significantly lower content was observed compared to the stressed treatments and ABA application seemed to have increased this situation (Table 3). Conversely, under 24 h of stress, the plants showed a different behavior; there was a noticeable decrease when compared to the other treatments. In relation to the increase of  $H_2O_2$  content following stress in the two maize hybrids, with ABA application, similar results were found by Kellos et al. (2008), where a tolerant maize lineage showed higher  $H_2O_2$  content when ABA was applied. The increase of  $H_2O_2$  at the beginning of stress can be explained by its role as a potent stress-signaling molecule (Møller et al. 2007). With increasing water stress,  $H_2O_2$  participation in the Haber–Weiss/Fenton reaction as a free radical attacking the cell membranes is noticeable (Queval et al. 2008). Exogenous treatment with ABA can increase internal ABA content (Table 2) and this can lead to an accumulation of  $H_2O_2$ , a signaling molecule for the activation of antioxidant enzymes. Such an explanation was previously demonstrated in maize (evaluating CAT activity and other enzymes; Jiang and Zhang 2002).

Drought stress causes a dramatic increase in reactive oxygen species (ROS) in plant cells. The accumulation of ROS causes membrane lipid peroxidation. MDA is a product of peroxidation of unsaturated fatty acids in phospholipids, and lipid peroxidation is responsible for cell membrane damage. ABA can cause the reduction of lipid peroxidation as expressed by MDA production in plant cells (Bhaskara et al.

2017; Kaya et al. 2018). The results showed that with the prolongation of drought stress, MDA content in the two varieties (ZD958 and XD20) accumulated greatly. Exogenous application of S-ABA led to a reduction in MDA content compared with no application, which was remarkable in a tolerant cultivar ( $p < 0.05$ ; Table 1). The latter may be related to the increase of antioxidant enzyme activity in ZD958 leaves and the better protection against oxidative stress. The current study confirmed that exogenous administration of S-ABA can alleviate damage caused by membrane lipid peroxidation in plants under water stress ( $p < 0.05$ ; Tables 3, 4). Our findings are in agreement with those of Ma et al. (2016), who show that the application of S-ABA reduces MDA content in wheat. The decrease of MDA and  $H_2O_2$  contents under PEG + S-ABA treatment, which can be due to the increase of antioxidant enzyme activity in leaves, promotes a better protection against oxidative stress (Moussa and Abdel-Aziz 2008).

Osmotic adjustment is an innate behavior of plants that helps to maintain water balance and adaptation to adverse conditions through the synthesis of different osmotic substances. The accumulation of osmotic substances under drought stress contributes to the regulation of plant physiological processes to adapt to adverse conditions. Generally, proline accumulation increases under stress conditions, which not only helps in maintaining cell turgor but also involves quenching-free radicals, maintaining sub-cellular structures, and buffering cellular redox potential. The current study showed that proline and soluble sugar accumulated rapidly in maize leaves after drought stress. Exogenous use of S-ABA hormone further increased the accumulation of proline and soluble sugar content, with ZD958 having a more significant response. Srivastava et al. (2009) showed that the application of exogenous ABA under water stress can increase the content of proline and soluble sugar in sugarcane seedlings ( $p < 0.05$ ) so, in maize, a similar response could be conducive to improving drought resistance. Under water stress, the use of S-ABA increases the content of proline and soluble sugar in maize leaves and reduces crop damage (Fig. 2).

A large number of experimental studies show that plants possess a well-defined antioxidant defense mechanism that eliminates hazardous free radicals (Moussa and Abdel-Aziz 2008; Suzuki and Katano 2018). A crop's protective enzymes (SOD, POD, CAT and APX) play an important role in the removal of peroxides and reactive oxygen species. Previous studies prove that higher enzyme activity or levels are important for inducing drought tolerance. The contribution of enzymatic antioxidants may ensure stress tolerance in plants subjected to long-term drought stress (De Souza et al. 2014; Gang et al. 2017). Jiang and Zhang (2002) reported that the application of exogenous ABA under stress conditions can induce the expression of related antioxidase

genes and increase the activity of related antioxidant defense enzymes. Plant antioxidant systems are induced and regulated by ABA signaling (Wang et al. 2016). In our study, the activity of antioxidant enzymes (including superoxide dismutase, catalase, peroxidase, ascorbate peroxidase, polyphenol oxidase, and phenylalanine ammonia-lyase) in maize leaves under drought stress increased rapidly ( $p < 0.05$ ). This was related to the rapid accumulation of endogenous ABA content under drought stress; however, the antioxidant enzyme activity decreased to different degrees after reaching a peak during different periods. An assumption was made that the equilibrium was destroyed due to the accumulation of large amounts of ROS. Our results are similar to the findings of Huang et al. (2013), who report that ABA spray causes an increase in related enzyme activity over that observed in untreated sugarcane. Exogenous administration of S-ABA compared with PEG treatment further increased the activity of antioxidant protective enzymes. Our results indicated that antioxidant enzyme content was higher in the tolerant cultivar than in the susceptible cultivar in all treatments ( $p < 0.05$ ; Figs. 3, 4). Anjum et al. (2017) demonstrate that the antioxidant response of maize under drought conditions is closely related to tolerance in maize varieties. Polyphenol oxidase (PPO), together with guaiacol peroxidase (POD), can act in diphenols and phenols to produce other phenolic compounds, such as flavonoids, which are important in scavenging free radicals (Pourcel et al. 2007). Whereas the Fazeli et al. (2007) study evidenced higher PPO activity in tolerant genotypes. Enhancement of L-phenylalanine ammonia lyase (PAL) activity was verified in drought stress. These data resemble those found by De Souza et al. (2014), where maize tolerant genotypes showed higher PAL activity. High PAL activity at the beginning of stress can also be important for a quick production of phenolic compounds, enhancing the protection against oxidative stress (Gholizadeh 2010; Hura et al. 2008).

The *Asr* gene family has drawn wide attention due to its excellent ability to relieve osmotic stress (Cakir et al. 2003). Much research indicates that after plants are subjected to stress (such as drought, low temperature, salt stress, etc.), the *Asr* gene responds to ABA regulation to alleviate any damage caused by adversity (Jia et al. 2016; Li et al. 2017; Shinozaki et al. 2003). Later studies reveal that the heterologous over-expression of plantain *Asr1* in *Arabidopsis* increases total soluble sugar levels in leaves (Dai et al. 2011). In addition, *Asr1* overexpression in maize plants increases yield and reduces sugar and amino acid levels; which suggests that *Asr1* also participates in plant growth through the regulation of this type of metabolite (Virilouvet et al. 2011). There is evidence suggesting that *Asr1* has a role in cell antioxidant activity. For instance, the overexpression of wheat *Asr1* in tobacco enhances the expression of ROS-related and stress-responsive genes under osmotic stress (Hu et al.

2013). Our results showed that the gene expressive level of *Asr1* in maize leaves increased under drought stress. It was speculated that the maize *Asr1* gene may play a role in signal transduction during drought resistance and then transmit more signals to induce downstream gene expression, thereby alleviating drought damage in maize ( $p < 0.05$ ). The expression of *Asr1* in exogenous S-ABA maize was significantly induced, and drought resistance was increased. This indicates that expression of the *Asr1* gene is regulated by S-ABA, which is consistent with the results of physiological indicators. Recently there have been similar reports for rice. With the application of exogenous S-ABA, the expression of the *Asr1* gene was significantly induced and the drought resistance of maize was increased ( $p < 0.05$ ; Fig. 5). This indicates that *Asr1* gene expression is regulated by S-ABA, and this is consistent with the physiological index results. Recently there have been similar reports in rice (Arenhart et al. 2016).

## Conclusion

In summary, under drought stress, maize became seriously damaged during the seed germination and the seedling stages. Seed soaking with 4 mg/kg S-ABA significantly enhanced the germination rate and mitigated the inhibitory effect of drought on seed germination. Foliar application of exogenous S-ABA increased endogenous ABA content, free proline, soluble sugar content, and *Asr1* level of gene expression; it also induced the enhancement of antioxidant enzyme activity (including SOD, CAT, POD, APX, PPO, and PAL), and reduced the accumulation of MDA, thereby improving drought resistance in maize. There were genotypic differences between the two maize varieties. Under drought stress, endogenous ABA content, osmotic substance content, antioxidant enzyme activity, and *Asr1* gene expression level were higher in the drought-resistant ZD958 variety. These results provide a basis for further study of the mechanism of S-ABA in mitigating damage caused by drought stress to maize seedlings; in the meantime they provide a theoretical reference for the screening of maize varieties.

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**Author Contributions** CY and FZ carried out the measurements, data analysis, and drafted the manuscript. XJ designed the experiments. XS, DS, and FH performed the experiment, XS, DS, FH, and XJ made substantial contributions to conception, and critically revised the

manuscript. All authors contributed to manuscript revision, read and approved the submitted version.

## Compliance with Ethical Standards

**Conflict of interest** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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